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SEQUENCE LISTING

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<120> METHODS AND COMPOSITIONS FOR
      DETERMINATION OF GLYCATED PROTEINS
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<140> 10/622,893
<141> 2003-07-17
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Thr Val Leu Asp Pro Tyr Pro Val Pro Ser Ala Ile Ser Ala Gly Asn
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Asp Val Asn Lys Val Ile Ser Ser Gly Gln Tyr Ser Asn Asn Lys Asp
Glu Ile Glu Val Asn Glu Ile Leu Ala Glu Glu Ala Phe Asn Gly Trp
Lys Asn Asp Pro Leu Phe Lys Pro Tyr Tyr His Asp Thr Gly Leu Leu
                                    90
Met Ser Ala Cys Ser Gln Glu Gly Leu Asp Arg Leu Gly Val Arg Val
                                105
Arg Pro Gly Glu Asp Pro Asn Leu Val Glu Leu Thr Arg Pro Glu Gln
                           120
       115
                                                125
Phe Arg Lys Leu Ala Pro Glu Gly Val Leu Gln Gly Asp Phe Pro Gly
                       135
Trp Lys Gly Tyr Phe Ala Arg Ser Gly Ala Gly Trp Ala His Ala Arg
                   150
                                        155
Asn Ala Leu Val Ala Ala Ala Arg Glu Ala Gln Arg Met Gly Val Lys
               165
                                    170
Phe Val Thr Gly Thr Pro Gln Gly Arg Val Val Thr Leu Ile Phe Glu
                                185
Asn Asn Asp Val Lys Gly Ala Val Thr Gly Asp Gly Lys Ile Trp Arg
                           200
Ala Glu Arg Thr Phe Leu Cys Ala Gly Ala Ser Ala Gly Gln Phe Leu
                       215
Asp Phe Lys Asn Gln Leu Arg Pro Thr Ala Trp Thr Leu Val His Ile
                                       235
Ala Leu Lys Pro Glu Glu Arg Ala Leu Tyr Lys Asn Ile Pro Val Ile
                245
                                    250
Phe Asn Ile Glu Arg Gly Phe Phe Phe Glu Pro Asp Glu Glu Arg Gly
                                265
Glu Ile Lys Ile Cys Asp Glu His Pro Gly Tyr Thr Asn Met Val Gln
                            280
Ser Ala Asp Gly Thr Met Met Ser Ile Pro Phe Glu Lys Thr Gln Ile
                        295
                                            300
Pro Lys Glu Ala Glu Thr Arg Val Arg Ala Leu Leu Lys Glu Thr Met
                    310
                                        315
Pro Gln Leu Ala Asp Arg Pro Phe Ser Phe Ala Arg Ile Cys Trp Cys
                325
                                    330
Ala Asp Thr Ala Asn Arg Glu Phe Leu Ile Asp Arg His Pro Gln Tyr
                                345
His Ser Leu Val Leu Gly Cys Gly Ala Ser Gly Arg Gly Phe Lys Tyr
                            360
Leu Pro Ser Ile Gly Asn Leu Ile Val Asp Ala Met Glu Gly Lys Val
                        375
                                            380
Pro Gln Lys Ile His Glu Leu Ile Lys Trp Asn Pro Asp Ile Ala Ala
                    390
                                        395
Asn Arg Asn Trp Arg Asp Thr Leu Gly Arg Phe Gly Gly Pro Asn Arg
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                                    410
Val Met Asp Phe His Asp Val Lys Glu Trp Thr Asn Val Gln Tyr Arg
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                               425
Asp Ile Ser Lys Leu
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```
Leu Gly Cys Gly Ala Ser Gly Arg Gly Phe Lys Tyr Leu Pro Ser Ile
                        375
                                            380
Gly Asn Leu Ile Val Asp Ala Met Glu Gly Lys Val Pro Gln Lys Ile
                    390
                                        395
His Glu Leu Ile Lys Trp Asn Pro Asp Ile Ala Ala Asn Arg Asn Trp
                405
                                    410
Arg Asp Thr Leu Gly Arg Phe Gly Gly Pro Asn Arg Val Met Asp Phe
                                425
His Asp Val Lys Glu Trp Thr Asn Val Gln Tyr Arg Asp Ile Ser Lys
                            440
Leu Lys Gly Glu Leu Glu Gly Leu Pro Ile Pro Asn Pro Leu Leu Arg
                        455
Thr Gly His His His His His
465
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tataccaacg ttaccgtgct ggacccctat cctgtcccta gcgccatctc cgccggaaac 180
gacgtgaaca aagtcattag cagtggccaa tattcgaata acaaagacga aatcgaagtg 240
aatgagatet tggeggaaga ggegtttaac ggttggaaga acgacceget tttcaaaccg 300
tattatcatg atacgggcct gctgatgtct gcttgctcgc aggagggcct ggatcgcctg 360
ggcgtccggg tacgtccggg cgaggatcct aatctggtgg aacttacccg cccggagcaa 420 '
tttcgtaaac tggccccgga aggcgtgttg caaggtgatt ttccgggttg gaaagggtac 480
tttqcqcqtt ccqqcqctqq ctqqqcacat gcaaggaatq ccttaqtqqc aqcaqcacqc 540
gaagcacage geatgggtgt aaaatttgtt actggcacce egeagggteg tgtagteaeg 600
ttaatetttg aaaataacga tgtaaaaggt gccgttacgg gcgatggcaa aatttggaga 660
geggaaegta catteetgtg tgetgggget agegegggte agtteetaga ttteaagaat 720
caacttcgac caaccgcttg gaccctggta cacattgcgt taaaaccgga agaacgtqcg 780
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gaggagcgcg gtgagattaa aatatgcgat gaacacccgg gctacacaaa tatqqtccaq 900
agtgcaqacq qcacqatqat qaqcattccg ttcqaaaaaa cccaqattcc aaaaqaaqcc 960
qaaacqcqcq ttcqqqccct qctqaaaqaq acaatqcccc aqctqqcaqa ccqtccattc 1020
agettegeae geatttgetg gtgtgeegat acegegaate gegaatteet gatagatega 1080
catccgcagt accacagtct tgtgttgggc tgtggtgcga gcggaagagg gtttaaatat 1140
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cacqaattaa tcaaqtqqaa cccqqacatt qcqqcqaacc qtaactqqcq tqatactctq 1260
qqqcqttttq qcqqtccaaa tcqtqtqatg gattttcatq atqtqaaqqa atqqaccaat 1320
gttcagtatc gtgatatttc caagctgaaa ggagagttgg aaggtaagcc aatccctaac 1380
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Gly Gly Gly Gly Gly Gly
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Gly Gly Gly Gly Gly Gly
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Ser Thr Glu Ser Ser Ile Ile Val Ile Gly Ala Gly Thr Trp Gly Cys
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20
                      25
Leu Leu Leu Leu Leu Leu
     35
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Ala Val Thr Lys Ser Ser Ser Leu Leu Ile Val Gly Ala Gly Thr Trp
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20
                       25
Thr Thr Thr Thr Thr Thr
     35
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  <211> 11
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  Cys Gln Asp Leu Pro Gly Asn Asp Asn Ser Thr
  <210> 14
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  Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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His His His His His
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Asp Thr Tyr Arg Tyr Ile
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Glu Tyr Met Pro Met Glu
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Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg
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Gln Tyr Pro Ala Leu Thr
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Gln Arg Gln Tyr Gly Asp Val Phe Lys Gly Asp
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